

- 1 -

piece 1, NC\_000913, ybiA\_dinG-, config: linear, direction: -, begin: 832322, end: 832154

Sequence alignment diagram showing the amino acid sequence of a protein across various species. The sequence starts with a methionine (Met) at position 1. It includes several leucine (Leu), phenylalanine (Phe), and tyrosine (Tyr) residues. A tryptophan (Trp) residue is present at position 10. A histidine (His) residue is also shown. The sequence ends with a valine (Val) residue. The alignment shows conservation of amino acids across the different species.

— നൂതനം ഉ മുൻസിപ്പാലിറ്റി

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{-----} sd-(9)-ir 832294 Gap 2.3 bits  
-----| sd-ir 832294 ybia_dinG- total 7.7 bits
```

p10 2.7 bi

2.4 bits  
6 bits

\*832160      \*

- 10 -